

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/721,553

Source: _____

Date Processed by STIC: _____

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 10/19/2004

PATENT APPLICATION: US/10/721,553

TIME: 08:58:09

Input Set : N:\Crf3\RULE60\10721553.raw.txt

Output Set: N:\CRF4\10192004\J721553.raw

```

1 <110> APPLICANT: Batra, Surinder K.
2   Hollingsworth, Michael A.
3   University of Nebraska Board of Regents
4 <120> TITLE OF INVENTION: Novel Gene That is Amplified and
5   Overexpressed in Cancer and Methods of Use Thereof
6 <130> FILE REFERENCE: UNMC63121
7 <140> CURRENT APPLICATION NUMBER: US/10/721,553
8 <141> CURRENT FILING DATE: 2003-11-25
9 <150> PRIOR APPLICATION NUMBER: US/09/647,143
10 <151> PRIOR FILING DATE: 2000-09-27
11 <150> PRIOR APPLICATION NUMBER: PCT/US99/06633
12 <151> PRIOR FILING DATE: 1999-03-26
13 <150> PRIOR APPLICATION NUMBER: 60/079,649
14 <151> PRIOR FILING DATE: 1998-03-27
15 <160> NUMBER OF SEQ ID NOS: 22
16 <170> SOFTWARE: FastSEQ for Windows Version 3.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 1937
20 <212> TYPE: DNA
21 <213> ORGANISM: Homo sapiens
22 <400> SEQUENCE: 1
23   ttctcgccccg cccacctcat ctcaaccac tttccgcggg gagcggcgcc aagctgggcc      60
24   ttctctggat caggcgctccc ctgaagtcgg cagcggccctc tgcgtccccc ttcggtccccg    120
25   ctaggacccc gtccgggctg ccgtcgccctc gtcgctatgg cgcccaccat ccagacccag      180
26   gccagcggg aggatggcca caggcccaat tcccaccgga ctctgcctga gaggtctgga      240
27   gtggtctgcc gagtcaagta ctgcaatagc ctccctgata tccccttcga ccccaagttc      300
28   atcacctacc ctttcgacca gaacaggttc gtccagtaca aagccacttc cttggagaaa      360
29   cagcaciaaac atgacctcct gactgagcca gacctggggg tcaccatcga tctcatcaat      420
30   cctgacacct accgcatcga cccaatggt cttctagatc cagctgatga gaaacttttg      480
31   gaagaggaga ttcaggcccc caccagctcc aagagatccc agcagcacgc gaaggtggtg      540
32   ccatggatgc gaaagacaga gtacatctcc actgagttca accgttatgg catctccaat      600
33   gagaagcctg aggtcaagat tggggtttct gtgaagcagc agttttaccga ggaagaaata      660
34   taaaaagaca gggatagcca gatcacagcc attgagaaga cttttgagga tgcccagaaa      720
35   tcaatctcac agcattacag caaaccgccg gtcacaccgg tggagggtcat gcctgtcttc      780
36   ccagacttta agatgtggat caatccatgt gctcagggtga tctttgactc agaccagacc      840
37   cccaaggaca cgagtgtgct agctgcgttg gagatgatgt ctcaggccat gattagggggc      900
38   atgatggatg aggaagggaa ccagtttgtg gcctatttcc tgctgtaga agagacgttg      960
39   aagaaacgaa agcgggacca ggaggaggag atggactatg caccagatga tgtgtatgac     1020
40   tacaaaattg ctcgggagta caactggaac gtgaagaaca aagctagcaa gggctatgag     1080
41   gaaaactact tcttcatctt ccgagagggt gacgggggtt actacaatga gttggaaacc     1140
42   aggggtccgcc ttagtaagcg cggggccaag gctgggggttc agtcaggcac caacgacctg     1200
43   cttgtggtca aacatcgga catgaatgag aaggaactgg aagctcagga ggcacggaag     1260
44   gccagctag aaaaccacga accggaggag gaagaggaag aggagatgga gacagaagag     1320

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45 aaagaagctg ggggctcaga tgaggagcag gagaagggca gcagcagtga gaaggagggc 1380
46 agtgaagatg agcactcggg cagcgagagt gaacgggagg aaggtgacag ggacgaggcc 1440
47 agtgacaaga gtggcagtgg tgaggacgag agcagcgagg atgaggcccg ggctgcccgt 1500
48 gacaaagagg agatcttttg cagtgatgct gattctgagg acgatgccga ctctgatgat 1560
49 gaggacagag gacaggccca aggtggcagt gacaatgatt cagacagcgg cagcaatggg 1620
50 ggtggccagc ggagccggag ccacagccgc agcgccagtc ccttccccag tggcagcgag 1680
51 cactcggccc aggaggatgg cagtgaagct gcagcttctg attccagtga agctgatagt 1740
52 gacagtgact gagtcccagg gcattcaggg ctgggttcaga caccattatt gtgagcagca 1800
53 aagcactttt ctagtggctt gtttgtgagc ctttcacttg tttgttcccc acccccaaac 1860
54 ctttgctggt aataaagtca acttctcttt aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 1920
55 aaaaaaaaaa aaaaaaa 1937

57 <210> SEQ ID NO: 2
58 <211> LENGTH: 531
59 <212> TYPE: PRT
60 <213> ORGANISM: Homo sapiens
61 <400> SEQUENCE: 2
62 Met Ala Pro Thr Ile Gln Thr Gln Ala Gln Arg Glu Asp Gly His Arg
63 1 5 10 15
64 Pro Asn Ser His Arg Thr Leu Pro Glu Arg Ser Gly Val Val Cys Arg
65 20 25 30
66 Val Lys Tyr Cys Asn Ser Leu Pro Asp Ile Pro Phe Asp Pro Lys Phe
67 35 40 45
68 Ile Thr Tyr Pro Phe Asp Gln Asn Arg Phe Val Gln Tyr Lys Ala Thr
69 50 55 60
70 Ser Leu Glu Lys Gln His Lys His Asp Leu Leu Thr Glu Pro Asp Leu
71 65 70 75 80
72 Gly Val Thr Ile Asp Leu Ile Asn Pro Asp Thr Tyr Arg Ile Asp Pro
73 85 90 95
74 Asn Val Leu Leu Asp Pro Ala Asp Glu Lys Leu Leu Glu Glu Ile
75 100 105 110
76 Gln Ala Pro Thr Ser Ser Lys Arg Ser Gln Gln His Ala Lys Val Val
77 115 120 125
78 Pro Trp Met Arg Lys Thr Glu Tyr Ile Ser Thr Glu Phe Asn Arg Tyr
79 130 135 140
80 Gly Ile Ser Asn Glu Lys Pro Glu Val Lys Ile Gly Val Ser Val Lys
81 145 150 155 160
82 Gln Gln Phe Thr Glu Glu Glu Ile Tyr Lys Asp Arg Asp Ser Gln Ile
83 165 170 175
84 Thr Ala Ile Glu Lys Thr Phe Glu Asp Ala Gln Lys Ser Ile Ser Gln
85 180 185 190
86 His Tyr Ser Lys Pro Arg Val Thr Pro Val Glu Val Met Pro Val Phe
87 195 200 205
88 Pro Asp Phe Lys Met Trp Ile Asn Pro Cys Ala Gln Val Ile Phe Asp
89 210 215 220
90 Ser Asp Pro Ala Pro Lys Asp Thr Ser Gly Ala Ala Ala Leu Glu Met
91 225 230 235 240
92 Met Ser Gln Ala Met Ile Arg Gly Met Met Asp Glu Glu Gly Asn Gln
93 245 250 255
94 Phe Val Ala Tyr Phe Leu Pro Val Glu Glu Thr Leu Lys Lys Arg Lys

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```

95          260          265          270
96 Arg Asp Gln Glu Glu Glu Met Asp Tyr Ala Pro Asp Asp Val Tyr Asp
97          275          280          285
98 Tyr Lys Ile Ala Arg Glu Tyr Asn Trp Asn Val Lys Asn Lys Ala Ser
99          290          295          300
100 Lys Gly Tyr Glu Glu Asn Tyr Phe Phe Ile Phe Arg Glu Gly Asp Gly
101          305          310          315          320
102 Val Tyr Tyr Asn Glu Leu Glu Thr Arg Val Arg Leu Ser Lys Arg Arg
103          325          330          335
104 Ala Lys Ala Gly Val Gln Ser Gly Thr Asn Ala Leu Leu Val Val Lys
105          340          345          350
106 His Arg Asp Met Asn Glu Lys Glu Leu Glu Ala Gln Glu Ala Arg Lys
107          355          360          365
108 Ala Gln Leu Glu Asn His Glu Pro Glu Glu Glu Glu Glu Glu Met
109          370          375          380
110 Glu Thr Glu Glu Lys Glu Ala Gly Gly Ser Asp Glu Glu Gln Glu Lys
111          385          390          395          400
112 Gly Ser Ser Ser Glu Lys Glu Gly Ser Glu Asp Glu His Ser Gly Ser
113          405          410          415
114 Glu Ser Glu Arg Glu Glu Gly Asp Arg Asp Glu Ala Ser Asp Lys Ser
115          420          425          430
116 Gly Ser Gly Glu Asp Glu Ser Ser Glu Asp Glu Ala Arg Ala Ala Arg
117          435          440          445
118 Asp Lys Glu Glu Ile Phe Gly Ser Asp Ala Asp Ser Glu Asp Asp Ala
119          450          455          460
120 Asp Ser Asp Asp Glu Asp Arg Gly Gln Ala Gln Gly Gly Ser Asp Asn
121          465          470          475          480
122 Asp Ser Asp Ser Gly Ser Asn Gly Gly Gly Gln Arg Ser Arg Ser His
123          485          490          495
124 Ser Arg Ser Ala Ser Pro Phe Pro Ser Gly Ser Glu His Ser Ala Gln
125          500          505          510
126 Glu Asp Gly Ser Glu Ala Ala Ala Ser Asp Ser Ser Glu Ala Asp Ser
127          515          520          525
128 Asp Ser Asp
129          530

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131 <210> SEQ ID NO: 3

132 <211> LENGTH: 20

133 <212> TYPE: DNA

134 <213> ORGANISM: Homo sapiens

135 <400> SEQUENCE: 3

136 agtgacaaga gtggcagtgg

20

138 <210> SEQ ID NO: 4

139 <211> LENGTH: 20

140 <212> TYPE: DNA

141 <213> ORGANISM: Homo sapiens

142 <400> SEQUENCE: 4

143 gaggacagag gacaggccca

20

145 <210> SEQ ID NO: 5

146 <211> LENGTH: 20

RAW SEQUENCE LISTING

DATE: 10/19/2004

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Input Set : N:\Crif3\RULE60\10721553.raw.txt

Output Set: N:\CRF4\10192004\J721553.raw

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147 <212> TYPE: DNA
148 <213> ORGANISM: Homo sapiens
149 <400> SEQUENCE: 5
150     cactcggccc aggaggatgg                                20
152 <210> SEQ ID NO: 6
153 <211> LENGTH: 20
154 <212> TYPE: DNA
155 <213> ORGANISM: Homo sapiens
156 <400> SEQUENCE: 6
157     gacagtgact gagtcccagg                                20
159 <210> SEQ ID NO: 7
160 <211> LENGTH: 19
161 <212> TYPE: DNA
162 <213> ORGANISM: Homo sapiens
163 <400> SEQUENCE: 7
164     ctggatggtg ggcgccata                                19
166 <210> SEQ ID NO: 8
167 <211> LENGTH: 19
168 <212> TYPE: DNA
169 <213> ORGANISM: Homo sapiens
170 <400> SEQUENCE: 8
171     cctgggtcccg ctttcgttt                                19
173 <210> SEQ ID NO: 9
174 <211> LENGTH: 19
175 <212> TYPE: DNA
176 <213> ORGANISM: Homo sapiens
177 <400> SEQUENCE: 9
178     ctaaggcgga ccctggttt                                19
180 <210> SEQ ID NO: 10
181 <211> LENGTH: 21
182 <212> TYPE: PRT
183 <213> ORGANISM: Homo sapiens
184 <400> SEQUENCE: 10
185     Arg Tyr Gly Ile Ser Asn Glu Lys Pro Glu Val Lys Ile Gly Val Ser
186     1           5           10           15
187     Val Lys Gln Gln Phe
188           20
190 <210> SEQ ID NO: 11
191 <211> LENGTH: 22
192 <212> TYPE: PRT
193 <213> ORGANISM: Homo sapiens
194 <400> SEQUENCE: 11
195     Glu Thr Arg Val Arg Leu Ser Lys Arg Arg Ala Lys Ala Gly Val Gln
196     1           5           10           15
197     Ser Gly Thr Asn Ala Leu
198           20
200 <210> SEQ ID NO: 12
201 <211> LENGTH: 18
202 <212> TYPE: DNA

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Output Set: N:\CRF4\10192004\J721553.raw

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203 <213> ORGANISM: Homo sapiens
204 <400> SEQUENCE: 12
205      ttcagtcagg caccaacg
207 <210> SEQ ID NO: 13
208 <211> LENGTH: 18
209 <212> TYPE: DNA
210 <213> ORGANISM: Homo sapiens
211 <400> SEQUENCE: 13
212      cgctggccac cccattg
214 <210> SEQ ID NO: 14
215 <211> LENGTH: 11
216 <212> TYPE: PRT
217 <213> ORGANISM: Homo sapiens
218 <400> SEQUENCE: 14
219      Arg Val Arg Leu Ser Lys Arg Arg Ala Lys Ala
220      1              5              10
222 <210> SEQ ID NO: 15
223 <211> LENGTH: 39
224 <212> TYPE: PRT
225 <213> ORGANISM: Drosophila melanogaster
226 <400> SEQUENCE: 15
227      Ser Asp Arg Arg Ser Asn Lys Pro Ile Met Glu Lys Arg Arg Arg Ala
228      1              5              10              15
229      Arg Ile Asn Asn Cys Leu Asn Glu Leu Lys Ala Asp Ile Leu Glu Lys
230      20              25              30
231      Thr Val Lys His Leu Gln Glu
232      35
234 <210> SEQ ID NO: 16
235 <211> LENGTH: 40
236 <212> TYPE: PRT
237 <213> ORGANISM: Homo sapiens
238 <400> SEQUENCE: 16
239      His Arg Pro Asn Ser His Arg Thr Leu Pro Glu Arg Ser Val Val Cys
240      1              5              10              15
241      Arg Val Lys Tyr Cys Asn Ser Leu Asp Ala Thr Ser Leu Glu Lys Gln
242      20              25              30
243      His Lys His Asp Leu Leu Thr Glu
244      35              40
246 <210> SEQ ID NO: 17
247 <211> LENGTH: 18
248 <212> TYPE: PRT
249 <213> ORGANISM: Homo sapiens
250 <400> SEQUENCE: 17
251      Ser Phe Gly Glu Leu Ala Leu Ile Tyr Gly Tyr Pro Arg Ala Ala Thr
252      1              5              10              15
253      Val Lys
255 <210> SEQ ID NO: 18
256 <211> LENGTH: 18
257 <212> TYPE: PRT

```

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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41,42
Seq#:1; Line(s) 43,44,45,46,47,48,49,50,51,52,53,54,55
Seq#:3; Line(s) 136
Seq#:4; Line(s) 143
Seq#:5; Line(s) 150
Seq#:6; Line(s) 157
Seq#:7; Line(s) 164
Seq#:8; Line(s) 171
Seq#:9; Line(s) 178
Seq#:12; Line(s) 205
Seq#:13; Line(s) 212

VERIFICATION SUMMARY

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